	OIPE
	CRF Erro Corrected by the STIC Systems Canch CRF Processing Date: 5/0/
V	umber: 07/07/,/32 Edited by:
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
i	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included: , '
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end o page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	eleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (enume to a Patentin bug). Sequences corrected:
	Other: conected spelling of Chlamydia globally

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

Input Set : A:\US08186269A.raw

								05			
	4	<110>		ANT: Bhatia				N.			
	5			, Yasir A.W	<i>N</i> .			•			
	6		Probst	, Peter							
	8										
	9										
	11										
C>	13	3 <140> CURRENT APPLICATION NUMBER: US/09/841,132									
	14	<141> CURRENT FILING DATE: 2001-04-23									
	16	<pre>< <160> NUMBER OF SEQ ID NOS: 599 < <170> SOFTWARE: FastSEQ for Windows Version 3.0/4.0 < <210> SEQ ID NO: 1</pre>									
	18										
		<211>									
		<212>	•								
		3 <213> ORGANISM: Chlamydia trachomatis									
		<400> SEQUENCE: 1									
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	27					accttcttta			120		
	28		,,	_		gtgaacgtat			180		
	29		_	-		atcattaaga			240		
	30					cgtaatatca			300		
						atgttccaaa			360		
	31								420		
	32				_	acgtgttcct	_		480		
	33.	_	cattet	ttttgttcgt	ttttgtgggt	attactgtat	Cilidadad	tatettagea			
	34	g 10105	ano to	NO 0					481		
			SEQ ID								
			LENGTH								
		<pre>< <212> TYPE: DNA </pre> <213> ORGANISM: Chlamydia trachomatis									
		. <400> SEQUENCE: 2 ! atcqttqqtq caqqacctat gcctcgcaca gagatcatta agaaaatgtg ggattacatt									
	42								60		
	43		-	-		aaacgtaata	_		120		
	44	gcta	aagttt	ttggaactga	aaaacctatc	gatatgttcc	aaatgacaaa	aatggtttct	180		
	45	caa							183		
			SEQ ID		•				•		
			LENGTH								
			TYPE:								
		0 <213> ORGANISM: Chlamydia trachomatis 2 <400> SEQUENCE: 3									
	52										
	53	gctg	cgacat	catgcgagct	tgcaaaccaa	catggacatc	tccaatttcc	ccttctaact	60		
	54	cgct	ctttgg	aactaatgct	gctaccgagt	caatcacaat	cacatcgacc		110		
	56	<210>	SEQ ID	NO: 4				•			
•	57	<211>	LENGTH	: 555							
	58	8 <212> TYPE: DNA 9 <213> ORGANISM: Chlamydia trachomatis									
	59										
		<400> SEQUENCE: 4									
	62				tatactactt	taagggaggc	ccttcgtatg	ccgcgcatca	60 -		
	63					taaaaataag			120		
			_		_	-					

Input Set : A:\US08186269A.raw

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64 tagggccage tetttetaaa gagattattg etagattgea gttgaateee gaagetagag
                                                                           180
65 ctgcagagtt gactgaggaa gaggttggtc gactaaacgc tcttttacag tcggattacg
                                                                           240
                                                                           300
66 ttgttgaagg ggatttgcgc cgtcgtgtgc aatctgatat caaacgtctg attactatcc
67 atgcttatcq tqqacaaaqa cataqacttt ctttqcctqt tcqtqqtcaq aqaacaaaaa
                                                                           360
68 caaattotog cacgogtaag ggtaaacgta aaactattgo aggtaagaag aaataataat
                                                                           420
   ttttaggaga gagtgttttg gttaaaaatc aagcgcaaaa aagaggcgta aaaagaaaac
                                                                           480
   aagtaaaaaa cattccttcg ggcgttgtcc atgttaaggc tacttttaat aatacaattg
                                                                           540
                                                                           555
  taaccataac agacc
73 <210> SEQ ID NO: 5
74 <211> LENGTH: 86
75 <212> TYPE: PRT
76 <213> ORGANISM: Chlamydia trachomatis
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79
80
    1
                                        10
   Ala Asp Leu Ala Ala Ile Val Gly Ala Gly Pro Met Pro Arg Thr Glu
81
82
                                    25
83
   Ile Ile Lys Lys Met Trp Asp Tyr Ile Lys Glu Asn Ser Leu Gln Asp
84
85
   Pro Thr Asn Lys Arg Asn Ile Asn Pro Asp Asp Lys Leu Ala Lys Val
86
                          . 55
87
   Phe Gly Thr Glu Lys Pro Ile Asp Met Phe Gln Met Thr Lys Met Val
                                             75
88
                        70
   Ser Gln His Ile Ile Lys
92 <210> SEQ ID NO: 6
93 <211> LENGTH: 61
94 <212> TYPE: PRT
95 <213> ORGANISM: Chlamydia trachomatis
97 <400> SEQUENCE: 6
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99
                                        10
    Trp Asp Tyr Ile Lys Glu Asn Ser Leu Gln Asp Pro Thr Asn Lys Arg
100
101
                 20
    Asn Ile Asn Pro Asp Asp Lys Leu Ala Lys Val Phe Gly Thr Glu Lys
102
103
                                 40
    Pro Ile Asp Met Phe Gln Met Thr Lys Met Val Ser Gln
107 <210> SEQ ID NO: 7
108 <211> LENGTH: 36
109 <212> TYPE: PRT
110 <213> ORGANISM: Chlamydia trachomatis
112 <400> SEQUENCE: 7
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114
                                          10
    Pro Leu Leu Thr Arg Ser Leu Glu Leu Met Leu Leu Pro Ser Gln Ser
115
                 20
116
117
     Gln Ser His Arg
118
             35
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Input Set : A:\US08186269A.raw

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120 <210> SEQ ID NO: 8
121 <211> LENGTH: 18
122 <212> TYPE: PRT
123 <213> ORGANISM: Chlamydia trachomatis
125 <400> SEQUENCE: 8
126 Leu Arg His His Ala Ser Leu Gln Thr Asn Met Asp Ile Ser Asn Phe
127
128 Pro Phe
131 <210> SEQ ID NO: 9
132 <211> LENGTH: 5
133 <212> TYPE: PRT
134 <213> ORGANISM: Chlamydia trachomatis
136 <400> SEQUENCE: 9
137 Leu Ala Leu Trp Asn
138
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140 <210> SEQ ID NO: 10
141 <211> LENGTH: 11
142 <212> TYPE: PRT
143 <213> ORGANISM: Chlamydia trachomatis
145 <400> SEQUENCE: 10
146 Cys Cys Tyr Arg Val Asn His Asn His Ile Asp
147
149 <210> SEQ ID NO: 11
150 <211> LENGTH: 36
151 <212> TYPE: PRT
152 <213> ORGANISM: Chlamydia trachomatis
154 <400> SEQUENCE: 11
155 Val Asp Val Ile Val Ile Asp Ser Val Ala Ala Leu Val Pro Lys Ser
156
                                         10
157
    Glu Leu Glu Gly Glu Ile Gly Asp Val His Val Gly Leu Gln Ala Arg
159
    Met Met Ser Gln
160
            35
162 <210> SEQ ID NO: 12
163 <211> LENGTH: 122
164 <212> TYPE: PRT
165 <213> ORGANISM: Chlamydia trachomatis
167 <400> SEQUENCE: 12
168 Met Pro Arg Ile Ile Gly Ile Asp Ile Pro Ala Lys Lys Lys Leu Lys
169
     1
     Ile Ser Leu Thr Tyr Ile Tyr Gly Ile Gly Pro Ala Leu Ser Lys Glu
170
171
     Ile Ile Ala Arg Leu Gln Leu Asn Pro Glu Ala Arg Ala Ala Glu Leu
172
173
                                 40
    Thr Glu Glu Glu Val Gly Arg Leu Asn Ala Leu Leu Gln Ser Asp Tyr
175
176
    Val Val Glu Gly Asp Leu Arg Arg Val Gln Ser Asp Ile Lys Arg
177
                         70
178 Leu Ile Thr Ile His Ala Tyr Arg Gly Gln Arg His Arg Leu Ser Leu
```

Input Set: A:\US08186269A.raw

```
179
                                                              95
     Pro Val Arg Gly Gln Arg Thr Lys Thr Asn Ser Arg Thr Arg Lys Gly
180
                 100
                                      105
     Lys Arg Lys Thr Ile Ala Gly Lys Lys
182
             115
183
185 <210> SEQ ID NO: 13
186 <211> LENGTH: 20
187 <212> TYPE: PRT
188 <213> ORGANISM: Chlamydia trachomatis
190 <400> SEQUENCE: 13
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192
      1
                                          10
193
    Val Phe Gly Thr
194
                 20
196 <210> SEQ ID NO: 14
197 <211> LENGTH: 20
198 <212> TYPE: PRT
199 <213> ORGANISM: Chlamydia trachomatis
201 <400> SEQUENCE: 14
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203
                                          10
      1
     Phe Gln Met Thr
204
205
                 20
207 <210> SEQ ID NO: 15
208 <211> LENGTH: 161
209 <212> TYPE: DNA
210 <213> ORGANISM: Chlamydia trachomatis
212 <400> SEQUENCE: 15
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                                                                             .60
213
                                                                             120
    ttacctacct cgcgacattc ggagctatcc gtccgattct gtttgtcaac aaaatgctgg
    cgcaaccgtt tctttcttcc caaactaaag caaatatggg a
                                                                             161
217 <210> SEQ ID NO: 16
218 <211> LENGTH: 897
219 <212> TYPE: DNA
220 <213> ORGANISM: Chlamydia trachomatis
222 <400> SEQUENCE: 16
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    acacagccca acaataaaat ggcaagggta gtaaataaga cgaagggaat ggataagact
                                                                             120
    attaaggttg ccaagtctgc tgccgaattg accgcaaata ttttggaaca agctggaggc
                                                                             180
225
     qcqqqctctt ccqcacacat tacaqcttcc caaqtqtcca aaqqattagg ggatgcgaga
                                                                             240
226
227
     actgttgtcg ctttagggaa tgcctttaac ggagcgttgc caggaacagt tcaaagtgcg
                                                                             300
228
    caaagcttct tctctcacat gaaagctgct agtcagaaaa cgcaagaagg ggatgagggg
                                                                             360
                                                                             420
229
    ctcacagcag atctttgtgt gtctcataag cgcagagcgg ctgcggctgt ctgtagcatc
230
     ateggaggaa ttacctacct egegacatte ggagetatee gteegattet gtttgtcaac
                                                                             480
    aaaatgctgg caaaaccgtt tctttcttcc caaactaaag caaatatggg atcttctgtt
                                                                             540
     agctatatta tggcggctaa ccatgcagcg tctgtggtgg gtgctggact cgctatcagt
                                                                             600
232
                                                                             660
233
    gcggaaagag cagattgcga agcccgctgc gctcgtattg cgagagaaga gtcgttactc
                                                                             720
234
     gaagtgccgg gagaggaaaa tgcttgcgag aagaaagtcg ctggagagaa agccaagacg
    ttcacqcqca tcaaqtatgc actcctcact atgctcgaga agtttttgga atgcgttgcc
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235
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Input Set : A:\US08186269A.raw

Output Set: C:\CRF3\05222001\I841132.raw

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236 gacgttttca aattggtgcc gctgcctatt acaatgggta ttcgtgcgat tgtggctgct
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                                                                             897
239 <210> SEQ ID NO: 17
240 <211> LENGTH: 298
241 <212> TYPE: PRT
242 <213> ORGANISM: Chlamydia trachomatis
244 <400> SEQUENCE: 17
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246
                                          10
247
     Lys Ala Phe Phe Thr Gln Pro Asn Asn Lys Met Ala Arg Val Val Asn
248
249
     Lys Thr Lys Gly Met Asp Lys Thr Ile Lys Val Ala Lys Ser Ala Ala
250
     Glu Leu Thr Ala Asn Ile Leu Glu Gln Ala Gly Gly Ala Gly Ser Ser
251
252
                              55
253
     Ala His Ile Thr Ala Ser Gln Val Ser Lys Gly Leu Gly Asp Ala Arg
254
                         70
255
     Thr Val Val Ala Leu Gly Asn Ala Phe Asn Gly Ala Leu Pro Gly Thr
256
257
     Val Gln Ser Ala Gln Ser Phe Phe Ser His Met Lys Ala Ala Ser Gln
258
                 100
                                      105
259
     Lys Thr Gln Glu Gly Asp Glu Gly Leu Thr Ala Asp Leu Cys Val Ser
                                  120
260
                                                      125
     His Lys Arg Arg Ala Ala Ala Val Cys Ser Ile Ile Gly Gly Ile
261
262
263
     Thr Tyr Leu Ala Thr Phe Gly Ala Ile Arg Pro Ile Leu Phe Val Asn
264
265
     Lys Met Leu Ala Lys Pro Phe Leu Ser Ser Gln Thr Lys Ala Asn Met
266
                     165
                                          170
267
     Gly Ser Ser Val Ser Tyr Ile Met Ala Ala Asn His Ala Ala Ser Val
268
269
     Val Gly Ala Gly Leu Ala Ile Ser Ala Glu Arg Ala Asp Cys Glu Ala
270
                                  200
271
     Arg Cys Ala Arg Ile Ala Arg Glu Glu Ser Leu Leu Glu Val Pro Gly
272
                             215
273
    Glu Glu Asn Ala Cys Glu Lys Lys Val Ala Gly Glu Lys Ala Lys Thr
274
                         230
                                              235
275
     Phe Thr Arg Ile Lys Tyr Ala Leu Leu Thr Met Leu Glu Lys Phe Leu
276
                                          250
     Glu Cys Val Ala Asp Val Phe Lys Leu Val Pro Leu Pro Ile Thr Met
277
278
279
     Gly Ile Arg Ala Ile Val Ala Ala Gly Cys Thr Phe Thr Ser Ala Ile
28.0
                                  280
                                                      285
             275
281
     Ile Gly Leu Cys Thr Phe Cys Ala Arg Ala
282
         290
                              295
284 <210> SEQ ID NO: 18
285 <211> LENGTH: 18
286 <212> TYPE: PRT
287 <213> ORGANISM: Chlamydia trachomatis
```

Please Note:

Use f n and/ r Xaa have been detected in the Sequence Listing. Pleas review th Sequence Listing t ensure that a c rresponding explanation is presented in the <220> to <223> fields of each sequenc which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/841,132

DATE: 05/22/2001 TIME: 16:14:06

Input Set : A:\US08186269A.raw

```
L:13 M:270 C: Current Application Number differs, Wrong Format
L:709 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:730 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46
L:758 \text{ M}:341 \text{ W}: (46) \text{ "n" or "Xaa" used, for SEQ ID$#:47}
L:1185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69
L:1265 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72
L{:}1266~M{:}341~W{:} (46) "n" or "Xaa" used, for SEQ ID#:72
L:1295 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73
L:1296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73
L:1297 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73
                     "n" or "Xaa" used, for SEQ ID#:76
L:1358 M:341 W: (46)
L:2310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:120
                     "n" or "Xaa" used, for SEQ ID#:175
L:3766 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:176
L:3970 M:341 W: (46)
                      "n" or "Xaa" used, for SEQ ID#:189
L:5193 M:341 W: (46)
L:5197 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:189
                     "n" or "Xaa" used, for SEQ ID#:189
L:5219 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:189
L:5221 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:189
L:5223 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:263
L:6854 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:264
L:6895 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:265
L:6931 M:341 W: (46)
                     "n" or "Xaa" used, for SEQ ID#:266
L:6972 M:341 W: (46)
L:7058 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:271
L:7316 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:287
L:17644 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:492
L:18675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:512
L:18677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:512
L:19520 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:524
L:19522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:524
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